TECH CENTER 1600/2900





P#330IPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/126,816A

DATE: 06/03/2002 TIME: 13:00:21

...,

Input Set : A:\98502bet.app

Output Set: N:\CRF3\06032002\I126816A.raw 3 <110> APPLICANT: VON EICHEL-STREIBER, CHRISTOPH BOQUET, PATRICE THELESTAM, MONICA 7 <120> TITLE OF INVENTION: METHOD OF INACTIVATION OF RAS SUBFAMILY PROTEINS AND AGENTS THEREFOR 10 <130> FILE REFERENCE: 98501/254992/bet 12 <140> CURRENT APPLICATION NUMBER: 09/126,816A C--> 13 <141> CURRENT FILING DATE: 2002-05-21 15 <150> PRIOR APPLICATION NUMBER: PCT/EP97/00426 16 <151> PRIOR FILING DATE: 1997-01-31 18 <150> PRIOR APPLICATION NUMBER: 96 101 469.3 19 <151> PRIOR FILING DATE: 1996-02-02 21 <160> NUMBER OF SEQ ID NOS: 6 23 <170> SOFTWARE: PatentIn Ver. 2.1 25 <210> SEQ ID NO: 1 26 <211> LENGTH: 31 27 <212> TYPE: DNA 28 <213> ORGANISM: Artificial Sequence 30 <220> FEATURE: 31 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer 33 <400> SEQUENCE: 1 31 34 ggggaatttt aatgagctca gttaacaaag c 37 <210> SEQ ID NO: 2 38 <211> LENGTH: 29 39 <212> TYPE: DNA 40 <213> ORGANISM: Artificial Sequence 42 <220> FEATURE: 43 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer 45 <400> SEQUENCE: 2 29 46 ttcagataat gtaggtacca agtctatag 49 <210> SEQ ID NO: 3 50 <211> LENGTH: 29 51 <212> TYPE: DNA 52 <213> ORGANISM: Artificial Sequence 54 <220> FEATURE:

55 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer

6/3/02

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66 <220> FEATURE:

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64 <213> ORGANISM: Artificial Sequence

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92 <212> TYPE: PRT
93 <213> ORGANISM: Clostridium sordellii
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97 1
                     5
                                        10
99 Phe Arg Ile Gln Glu Asp Glu Tyr Val Ala Ile Leu Asn Ala Leu Glu
               . 20
                                     25
102 Glu Tyr His Asn Met Ser Glu Ser Ser Val Val Glu Lys Tyr Leu Lys
105 Leu Lys Asp Ile Asn Asn Leu Thr Asp Asn Tyr Leu Asn Thr Tyr Lys
                             55
108 Lys Ser Gly Arg Asn Lys Ala Leu Lys Lys Phe Lys Glu Tyr Leu Thr
                         70
                                             75
111 Met Glu Val Leu Glu Leu Lys Asn Asn Ser Leu Thr Pro Val Glu Lys
                     85
                                         90
114 Asn Leu His Phe Ile Trp Ile Gly Gly Gln Ile Asn Asp Thr Ala Ile
                100
                                    105
115
117 Asn Tyr Ile Asn Gln Trp Lys Asp Val Asn Ser Asp Tyr Thr Val Lys
118
            115
                                120
120 Phe Val Tyr Asp Ser Asn Ala Phe Leu Ile Asn Thr Leu Lys Lys Thr
                                                140
                            135
123 Ile Val Glu Ser Ala Thr Asn Asn Thr Leu Glu Ser Phe Arg Glu Asn
                        150
                                            155
126 Leu Asn Asp Pro Glu Phe Asp Tyr Asn Lys Phe Tyr Arg Lys Arg Met
                    165
                                        170
129 Glu Ile Ile Tyr Asp Lys Gln Lys His Phe Ile Asp Tyr Tyr Lys Ser
                                    185
                                                         190
130
                180
132 Gln Ile Glu Glu Asn Pro Glu Phe Ile Ile Asp Asn Ile Ile Lys Thr
                                200
135 Tyr Leu Ser Asn Glu Tyr Ser Lys Asp Leu Glu Ala Leu Asn Lys Tyr
                            215
138 Ile Glu Glu Ser Leu Asn Lys Ile Thr Ala Asn Asn Gly Asn Asp Ile
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	_		225	240
130 225	230		235	Len Tvr Asn
139 225 141 Arg Asn Leu Gl	u Lys Phe Al	a Asp Glu Asp	Leu var mra	255
142	245	250	l . wla wla Ger	Asp Ile Leu
142 144 Gln Glu Leu Va	l Glu Arg Tr	p Asn Leu Ala	l Ald Ald Sci	270
145 26	0	265	val mor Lau	Asp Val Asp
145 26 147 Arg Ile Ser Me	t Leu Lys Gl	u Asp Gly Gly	Val Tyl ned	C ASP TWO
14/ AIG 110 201		280	20.	Nan Lvs Pro
148 275 150 Ile Leu Pro G	v Ile Gln Pr	co Asp Leu Pho	E Lys Ser II	ASH LIB 120
150 He Bed 110 0	29	95	300	. clu Ala Tle
151 290 153 Asp Ser Ile T	or Asn Thr Se	er Trp Glu Me	t Ile Lys Lei	320
153 ASP Set TIE T	310		315	- 3-m Pho ASD
154 305 156 Met Lys Tyr L	ve Glu Tvr I	le Pro Gly Ty	r Thr Ser Ly	S ASII PIIE ASP
156 Met Lys Tyr D	325	33	0	333
157 159 Met Leu Asp G	ın Glu Val G	ln Arg Ser Ph	e Glu Ser Al	a Leu Ser Ser
159 Met Leu Asp G	AO	345		350
160 3 162 Lys Ser Asp L	40 Cor Clu T	le Phe Leu Pr	o Leu Asp As	p Ile Lys var
162 Lys Ser Asp L	ys ser dia i	360	. 36	5
163 355 165 Ser Pro Leu G	ı vel tvc T	le Ala Phe Al	a Asn Asn Se	r Val Ile Asn
165 Ser Pro Leu G	ilu vai nys i	75	380	
166 370 168 Gln Ala Leu I		ve Asp Ser Ty	r Cys Ser As	sp Leu Val IIe
168 Gln Ala Leu I	Te ser Leu L	AP COT -	395	400
169 385 171 Asn Gln Ile I	390	our Ivs Ile Le	eu Asn Asp As	sn Leu Asn Pro
171 Asn Gln Ile I	Lys Asn Arg 1	1y1 Ly3 110 - 4:	10	415
172	405	san Dhe Asn T	hr Thr Met Ly	s Ile Phe Ser
172 174 Ser Ile Asn (	Glu Gly Thr F	425		430
175 177 Asp Lys Leu	420	a-m Aan Glu A	sp Asn Met M	et Phe Met Ile
177 Asp Lys Leu	Ala Ser IIe S	440	4	45
178 435 180 Lys Ile Thr	_	440 val Clv D	he Ala Pro A	sp Val Arg Ser
180 Lys Ile Thr	Asn Tyr Leu	TAS AUT GIA I	460	
181 450 183 Thr Ile Asn		455	or Thr Gly A	la Tyr Gln Asp
183 Thr Ile Asn	Leu Ser Gly	bro gry var r	475	480
184 465 186 Leu Leu Met	470	- a mbr 7	en Tle His I	eu Leu Glu Pro
186 Leu Leu Met	Phe Lys Asp	Asn Ser Till F	1911 110 11-1 -	495
187	485	T	the Lys Ile S	er Gln Leu Thr
187 189 Glu Leu Arg	Asn Phe Glu	Phe Pro Lys	III LYS IIC -	510
190	500	505	oho Aen Gln A	la Arg Ala Lys
192 Glu Gln Glu	Ile Thr Ser	Leu Trp Ser	Pile Asir Ori	525
193 515 195 Ser Gln Phe		520	ner Dho Glu (	lv Ala Leu Gly
195 Ser Gln Phe	Glu Glu Tyr	Lys Lys GIY	ryr Phe Gra (	, <b>-</b> 1
196 530 198 Glu Asp Asp		535	J40 mb~ Val l	en Asp Lys Asp
108 Glu ASD ASP	Asn Leu Asp	Phe Ala Gin	ASH THE VOL	560
199 545	550			Arg Asn Lvs Glu
201 Tur Val Ser	Lys Lys Ile	Leu Ser Ser	Met Lys Thi	Arg Asn Lys Glu 575
201 171 (41 001	565		570	TIA SAT TVT Glu
204 mur Tle His	Tvr Ile Val	Gln Leu Gln	GIY ASP LYS	Ile Ser Tyr Glu 590
204 191 110 1110	580	585		cor Tle Len Tyr
200 Ala Car CVS	Asn Leu Phe	Ser Lys Asp	Pro Tyr Ser	Ser Ile Leu Tyr 605
207 Ald Sel Cys		600	<del>_</del>	TUD TUR TIAL ALA
200	Tle Glu Glv	Ser Glu Thr	Ala Tyr Tyr	Tyr Tyr Val Ala
	. 110 010 -01	615	620	
211 610				

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213 Asp	Ala G	lu Il	e Ly	rs G	lu I 30	le A	Asp	Lys	Tyr	Arg	ı Il	e P	ro :	ľyr	Gln	116 64	<b>≘</b> 0
214 625 216 Ser	Asn L	ys Ar	g As	n I	le I	ys :	Leu	Thr	Phe	Ile	e Gl	у Н	is (	Gly	Lys 655	Se	r
217 219 Glu	Phe A	sn Th	o4 ar As	sp T	hr I	he	Ala	Asn	Leu	Ası	y Va	1 A	sp	Ser 670	Leu	Se	r
220 222 Ser	Glu I	be [le G]	ou Lu Tl	hr I	le l	Ĺeu	Asn	Leu	Ala	Ly	s Al	.a A	sp 585	Ile	Ser	Pr	0
223 225 Lys	Tyr 1	575 [le G]	lu I	le A	sn :	Leu	Leu	Gly	Cys	As	n Me	et E	he	Ser	Tyr	Se	r
226 228 Ile	690 Tyr <i>P</i>	Ala G	lu G	lu I	hr	695 Tyr	Pro	Gly	Lys	ь Le 71	u Le	eu I	Leu	Lys	Ile	Ьў 72	rs 20
229 705 231 Asp	Arg '	Val S	er G	lu I	Leu	Met	Pro	Ser	: Ile	e Se	r G	ln A	Asp	Ser	11e 735	Th	ır
232 234 Val	Ser i	Ala A	sn G	25 1n :	Гуr	Glu	Val	Arg	Il	e As	n G	lu (	Glu	Gly 750	Lys	A 1	rg
235 237 Glu	Ile	7 Leu A	40 .sp H	lis	Ser	Gly	Lys	Tr	, Il	e As	n L	ys	Glu 765	Glu	Ser	: I.	le
238 240 Ile	Lys	755 Asp I	le S	Ser	Ser	Lys	Glu	ту	r Il	e Se	er P	he 80	Asn	Pro	Lys	s G	lu
241 243 Asn	770 Lys	Ile 1	le V	/al	Lys	775 Ser	Lys	з Ту	r Le	u Hi	is G	lu	Leu	Ser	Thi	r L 8	eu 00
244 785 246 Leu	Gln	Glu I	[le ]	Arg	790 Asn	Asn	Ala	a As	n Se	r Se	er A	sp	Ile	Asp	Let 81	ı G 5	lu
247 249 Lys	T.VS	Val 1	Met :	805 Leu	Thr	Glu	Су	s Gl	81 u II	.0 _e A	sn V	al	Ala	Sei	r As:	n I	le
250 252 Asp	) Lys	Gln	820 Tle	Val	Glu	Gly	Ar	82 g Il	5 e G	Lu G	lu A	Ala	Lys	Ası	n Le	u I	hr
252 ASE 253 255 Sei	, Arg	835 Ser	Tle	Asn	Tvr	Ile	84 Ly	0 s As	n G	lu P	he 1	Lys	Leu	Il	e Gl	u S	Ser
255 Sei 256 258 Ile	850	361	Cor	LOU	−∡ ∵Ф∨г	855 Ast	5 Le	u · Ly	s H	is G	ln i	860 Asn	Gly	, Le	u As	p P	Asp
258 Ile 259 86 261 Se	e Ser 5	Asp	ser	Leu	870	Cl	1 Ac	n T	le S	8 er I	75 Jys '	Thr	Glı	ı As	n Gl	y 1	380 Phe
261 Se: 262 264 Ar	r His	Phe	He	885	PHE	. 61	_ (1	. T	8 ar G	90 1 v 7	.sn	Ser	Ile	e Ph	89 e Il	15 .e (	Glu
264 Ar	g Ile	Arg	Phe 900	Ile	Asn	т г	S G1	9	05	- y -	Dhr	uic	т1.	91 - Se	.0 er Lv	7S	Glu
265 267 Th	r Glu	Lys 915	Glu	Ile	Phe	e Se	r GJ 92	20 T	у г	.1a .		11-0	92	5 n G1	v I.	/S	Leu
268 270 Il	e Ser	Asn	Ile	Lys	Ası	Th 93	r II 5	Le P	ne A	sp A	-2	940	. AS	n mi	or T.	211	Asn
271 273 Va	l Lys	Lys	Val	Asn	Lei 95	u As O	p A	la A	la H	Iis (	GLu 955	vaı	AS	11 11	11 D		960
274 94 276 Se	r Ala	n Phe	Phe	11e 965	G1:	n Se	r L	eu I	le (	31u 970	Tyr	Asr	1 Th	r Tl	nr L	y 5 75	g_u
277 279 Se	er Lei	ı Ser	Asn	Leu	ı Se	r Va	ıl A	la M	et 1 85	Lys	Val	Glr	n Va	1 T; 9	yr A 90	la	GIN
280 282 Le	eu Phe	e Ser	980 Thr	Gly	, Le	u As	n T	hr I	le '	Fhr	Asp	Ala	a Se	r L	ys V	al	Val
283 285 G	lu Le	995 u Val	Ser	Th	r Al	a Le	eu A	sp (	lu '	Thr	Ile	As	p Le	eu L	eu P	ro	Thr

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286 1010 1015 1020 288 Leu Ser Glu Gly Leu Pro Ile Ile Ala Thr Ile Ile Asp Gly Val Ser 1030 1035
288 Leu Ser Glu Gly Hed 110 110 1035 1040
289 1025 1030 1033 289 1025 1030 1033 291 Leu Gly Ala Ala Ile Lys Glu Leu Ser Glu Thr Asn Asp Pro Leu Leu 291 Leu Gly Ala Ala Ile Lys Glu Leu Ser Glu Thr Asn Asp Pro Leu Leu
291 Leu Gly Ala Ala 215 21- 1050 1055
292 1045 1050 294 Arg Gln Glu Ile Glu Ala Lys Ile Gly Ile Met Ala Val Asn Leu Thr 294 Arg Gln Glu Ile Glu Ala Lys Ile Gly Ile Met Ala Val Asn Leu Thr
294 Arg Gin Giu ile Giu Aiu 275 1065 1070
295 The Ale The Val Thr Ser Ala Leu Gly Ile Ala Ser Gly
295 1060 1065 297 Ala Ala Ser Thr Ala Ile Val Thr Ser Ala Leu Gly Ile Ala Ser Gly 297 Ala Ala Ser Thr Ala Ile Val Thr Ser Ala Leu Gly Ile Ala Ser Gly 1085
298 1075 1080 208 209 209 209 209 209 209 209 209 209 209
300 Phe Ser Ile Leu Leu Val Pio Hea Mia 327
301 1090 1095 1100 303 Ser Leu Val Asn Asn Glu Leu Ile Leu Gln Asp Lys Ala Thr Lys Val 1110 1115
303 Ser Leu Val Asn Asn Glu Leu 11e Beu Gli 115 1120
304 1105 1110 1113 Thr Glu Gly Ala Phe
304 1105 1110 1113 306 Ile Asp Tyr Phe Lys His Ile Ser Leu Ala Glu Thr Glu Gly Ala Phe 1135 1130
307 1125 1125 Not Pro Gln Asp Asp Leu Val Leu
200 Thr Leu Leu Asp Asp Lys IIe IIe Met FIO GIA 1150
309 Thr Leu Leu Asp Asp Lys 110 1145 1150 1140 1140 1145 1150
310 1140 1145 310 Ser Glu Ile Asp Phe Asn Asn Ser Ile Thr Leu Gly Lys Cys Glu 312 Ser Glu Ile Asp Phe Asn Asn Ser Ile Thr Leu Gly Lys Cys Glu
313 1155 1160 1163 1163 313 1155 1160 1160 1163
215 The Trn Arg Ala Glu Gly Gly Ser Gly HIS INT 100
316 1170 1175 1180 1180 316 Thr Tyr Arg Lys Pro Trp Leu
The pho pho ser ser pro ser lie in 111 mg = 1
318 Asp His Phe Phe Ser Ser Flo Ser 1200 1195 1200
319 1185 1190 1193 321 Ser Ile Tyr Asp Val Leu Asn Ile Lys Lys Glu Lys Ile Asp Phe Ser 1210 1215
321 Ser 11e 1yl Asp val 205 1210 1215
322 1205 1210 324 Lys Asp Leu Met Val Leu Pro Asn Ala Pro Asn Arg Val Phe Gly Tyr 1225 1230
324 Lys Asp Let 1220 1225 1230
325 1220 1225 327 Glu Met Gly Trp Thr Pro Gly Phe Arg Ser Leu Asp Asn Asp Gly Thr
327 Glu Met Gly TIP III FIO GI
328 1235 and I have the Arg Asp His Tyr Glu Gly Gln Phe Tyr Trp
330 Lys Leu Leu Asp Arg 11e Arg Asp 1260 1255 1260
331 1250 1255 Ala Leu Ile Thr Lys Leu Lys Pro
331 1250 1255 1260 333 Arg Tyr Phe Ala Phe Ile Ala Asp Ala Leu Ile Thr Lys Leu Lys Pro 1270 1275 1280
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334 1265 1270 1275 336 Arg Tyr Glu Asp Thr Asn Val Arg Ile Asn Leu Asp Gly Asn Thr Arg 1295 1290 1295
337 1285 The Thr Glu Glu Ile Arg Lys Asn Leu
337 1285 1290 339 Ser Phe Ile Val Pro Val Ile Thr Thr Glu Gln Ile Arg Lys Asn Leu 1300 1305
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342 Ser Tyr Ser Phe Tyr Gly Ser Gly Gly Sel Tyl Sel 1325
342 Sel Tyl 325 343 1315 1320 1325 345 Pro Tyr Asn Met Asn Ile Asp Leu Asn Leu Val Glu Asn Asp Thr Trp 1335 1340
345 Pro Tyr Asn Met Asn Ile Asp Leu Asn Leu var Gid Hen 451
346 1330 1335 The Thr Ile Glu Ser Asp
346 1330 1335 348 Val Ile Asp Val Asp Asn Val Lys Asn Ile Thr Ile Glu Ser Asp 1350 1355
349 1345 1350 1350 1350 1350 Ser Lys Leu Asn
349 1345 1350 1350 Leu Asn 351 Glu Ile Gln Lys Gly Glu Leu Ile Glu Asn 11e Leu Ser Lys Leu Asn 1370 1375
1370 1365 1370 The Tio Asp Phe Tyr
352 1365 1370 354 Ile Glu Asp Asn Lys Ile Ile Leu Asn Asn His Thr Ile Asn Phe Tyr 1385 1390
1385 1380 1385 1390
355 1380 1363 357 Gly Asp Ile Asn Glu Ser Asn Arg Phe Ile Ser Leu Thr Phe Ser Ile
358 1395

VERIFICATION SUMMARY

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date